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Procedures for the Semi-Automated Extraction of DNA

1 Scope

These procedures apply to DNA personnel performing semi-automated extraction and purification of deoxyribonucleic acid (DNA) using the QIAcube® and/or EZ1® Advanced XL (EZ1®) in the DNA Casework Unit (DCU) or Scientific and Biometrics Analysis Unit (SBAU) and DNA personnel that perform the associated quality control procedures.

2 Equipment/Materials/Reagents

Equipment/Materials

- General laboratory supplies (e.g., tubes, pipettes, vortex, centrifuge)
- Incubator (Thermo MaxQ 4450 or 4000, Thermo 6841, Labline Imperial III, Heratherm IGS 100, or equivalent) or thermomixer (Eppendorf Thermomixer 5350s or equivalent)
- Costar® spin baskets, or equivalent (for differential extractions)
- Qiagen[®] Lyse & Spin Baskets and Collection Tubes, or equivalent (*for normal extractions*)
- QIAcube®
- QIAcube[®] consumables (e.g., 1000 μL wide-bore filter-tips, reagent bottles, rotor adapters, QIAcube[®]-compatible 1.5 mL microcentrifuge tubes)
- EZ1® Advanced XL
- EZ1® DNA Investigator Kit (e.g., filter tips, tip holders, elution tubes, reagent cartridges)

Reagents

- Buffer ATL (for EZ1 female fraction)
- Buffer G2 (for EZ1 normal lysis and EZ1 male fraction)
- 1M Dithiothreitol (DTT)
- Proteinase K
- Reagent Grade Water

3 Standards and Controls

At least one extraction control (i.e., reagent blank) must be processed in parallel with each extraction batch. The reagent blank(s) will be processed as the last sample(s) in the batch.

For evaluation of the extraction controls, refer to the appropriate interpretation procedure of the *DNA Procedures Manual* (i.e., DNA 233, DNA 215, DNA 410).

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4 Procedures

Refer to the DNA Procedures Introduction (i.e., DNAQA 600) for applicable general precautions and cleaning instructions.

Ensure the appropriate fields (i.e., instruments, reagents) in the Sample Tracking and Control Software (STACS) are completed, as necessary.

4.1 Normal Lysis

The samples should be in Lyse & Spin baskets in corresponding tubes when using an incubator. If using a thermomixer, the Lyse & Spin baskets should not be used.

4.1.1	Create master mix using the recipe below.	
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Normal Lysis Master Mix

Reagent	μL per sample
Buffer G2	470
Pro K	15
DTT	15

4.1.2	Add 450 μL master mix.	
4.1.3	Incubate the tubes in an incubator at 56°C with agitation (generally 200 rpm) for ~1 hour.	
4.1.4	Spin the tubes (generally between 9,000 and 13,000 rpm for 5 minutes). Discard the basket.	

If the lysate does not completely flow through the basket, additional spins may be added. If necessary, lysate remaining in the basket may be manually transferred to the sample tube. Additional manipulations will be recorded in the case notes. If the volume in the sample tube is significantly different than the expected volume a case note will be made.

4.1.5	Process the lysates on the EZ1® following the steps in section 4.3.	
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4.2 Differential Lysis and QIAcube® Fractionation

The samples should be in QIAcube® compatible tubes. Lyse & Spin baskets must NOT be used for differential extractions.

4.2.1	Create the epithelial (F) fraction master mix.	
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If precipitate has formed in the Buffer ATL, heat it, generally at 56°C, until precipitate is no longer visible.

Female Fraction Lysis Master Mix

Reagent	μL per sample
Buffer ATL	160
Reagent Grade Water	320
Pro K	20

4.2.2	Add 450 μL master mix.	
4.2.3	Incubate the tubes at 56°C with agitation (generally in a thermomixer at 900 rpm) for ~1.5 hours.	
4.2.4	If necessary, quick spin and transfer the substrate to a basket. Spin the tubes (generally between 9,000 and 13,000 rpm for 5 minutes). Discard the basket	
	Discard the basket.	
4.2.5	Ensure consumables, reagent grade water, and lysate tubes are properly loaded onto the OIAcube® instrument.	

Consumables include: bottles containing reagent grade water, $1000~\mu L$ wide-bore tips, rotor adapters, and F fraction collection tubes. A shaker rack plug must be in place next to each F fraction collection tube.

Appendix A has additional guidance for loading the QIAcube®.

4.2.6	Initiate the "Separate and Lyse 12A Mod" protocol on the QIAcube [®] .	

The bottle in position 1 will contain reagent grade water, not Buffer G2 as prompted. Be aware that lysates are 450 μ L, not 500 μ L as referenced in the QIAcube[®] prompt.

4.2.7	At the completion of the run, tubes containing the F fraction should be	
	removed and capped.	
	Refill consumables and reagent grade water as necessary.	

The M fractions continue processing on the QIAcube®. Processing of the F fractions resumes at

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section 4.3, either independently or with the M fraction lysates.

4.2.8	Initiate the "Separate and Lyse 12B Mod" protocol on the QIAcube®.	
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The bottles will contain reagent grade water, not Buffer G2 as prompted.

4.2.9	At the completion of the run, remove and close the tubes containing the M	
	fraction(s).	

The M fraction tubes should contain approximately 50 µL of liquid following this run. If excess liquid remains, the tubes may be centrifuged (generally between 9,000 and 13,000 rpm) for 5 minutes and the excess supernatant manually removed and discarded while avoiding the pellet. If this step is necessary, a case note will be made.

4.2.10	A slide for microscopy may be prepared from the M fraction according to	
	Serology procedure for Christmas tree staining (i.e., Sero 112).	

4.2.11 Ensure the M fraction master mix is created.

Male Fraction Lysis Master Mix

Reagent	μL per sample
Buffer G2	375
Pro K	24
DTT	94

4.2.12	Add 450 μL master mix to each M fraction tube.	
4.2.13	Vortex, quick spin, and incubate the tubes at 70°C with agitation (generally in	
	a thermomixer at 900 rpm) for ~10 minutes.	
4.2.14	Vortex the tubes vigorously (~10 seconds) and quick spin.	
4.2.15	Process the lysates on the EZ1® following the steps in section 4.3.	

Ensure waste is disposed of properly.

The water bottles on the QIAcube® should be emptied at the end of each day of use.

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4.3 Processing Lysates on the EZ1® Advanced XL

Lysates may be stored refrigerated for up to 6 days prior to processing on the EZ1[®]. Ensure that no precipitate remains in the lysates prior to processing.

	Ensure the EZ1® is UV irradiated for 20 minutes prior to initial use each day.	
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T.J.1	Libert the L21 15 0 v irradiated for 20 minutes prior to minute disc each day.	

If necessary, transfer the lysate to an $EZ1^{\mathbb{R}}$ compatible tube.

4.3.3 Ensure consumables, lysate tubes, and barcoded elution tubes are properly loaded onto the EZ1[®]. See Figure 1.

If precipitate has formed in the EZ1® cartridges, heat them, generally at 56°C, until precipitate is no longer visible.

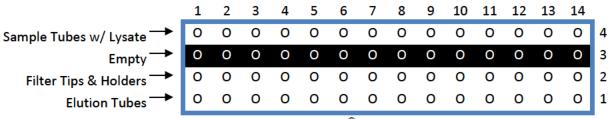


Figure 1 - Loading the EZ1® Advanced XL

4.3.4 Ensure the Large Volume protocol with elution into 50 μL of water has been selected and start the EZ1[®].

Disregard the reference to MTL Buffer in the prompts on the EZ1®.

4.3.5 At the completion of the run, remove and cap elution tubes.

If the volume in the elution tube is significantly different than the expected volume a case note will be made.

4.3.6 If necessary, combine appropriate extracts.

If combined samples will be concentrated using the Speed-Vac or Vacufuge, a maximum of eight 50 µL extracts may be combined into one tube.

4.3.7 Ensure waste is disposed of properly.
Ensure the piercing units on the EZ1® are cleaned after daily use and clean the tray, worktable, and racks, if needed.

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Bleach should not be used on the QIAcube® or the EZ1® Advanced XL. 70% ethanol should be used to clean instrument surfaces.

4.4 Combining/Concentrating Extracted Samples Using a Microcon Filter

For concentrating samples using the Speed-Vac or Vacufuge, refer to the procedures for quantification of nuclear DNA (i.e., DNA 226).

4.4.1	Vortex and quick spin the extract tubes. Transfer the extract for each sample	
	being combined/concentrated into a labeled microcon assembly.	

NOTE: Corresponding reagent blanks must also be microconned.

4.4.2	Spin the tubes to draw the fluid through the membrane (generally between	
	6,000 and 8,000 rpm for 10 minutes).	

Speed and time may be increased to draw fluid through membrane, but to avoid damage to the membrane, excess speed and time should not be used.

If additional spins do not reduce the volume, the affected sample(s) may continue with processing at step 4.6.4. Record the final volume.

4.4.3	Add reagent grade water (generally 15 µL).	
4.4.4	Invert microcon into a new, labeled tube.	
4.4.5	Spin the tubes (generally between 9,000 and 13,000 rpm for 5 minutes).	

Ensure the final tubes are barcoded.

5 Sampling

Not applicable.

6 Calculations

Not applicable.

7 Measurement Uncertainty

Not applicable.

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8 Limitations

The quantity and quality of the DNA present within any biological material ultimately determines if a DNA extraction is successful.

9 Safety

- 9.1 All evidence containing or contaminated with blood or other potentially infectious materials will be considered infectious regardless of the perceived status of the source individual or the age of the material. Follow the "Safe Work Practices and Procedures," "Bloodborne Pathogen (BBP) Exposure Control Plan (ECP)," "Personal Protective Equipment Policy," and "Chemical Hygiene Plan" sections of the *FBI Laboratory Safety Manual*.
- **9.2** Refer to the "Hazardous Waste Disposal" section of the *FBI Laboratory Safety Manual* for important information concerning proper disposal of the chemicals used in these procedures as well as the biohazardous wastes generated.
- **9.3** Procedural Specific Chemical Hazards:
 - Solutions of Proteinase K can be irritating to mucous membranes. Use eye protection when handling.
 - EZ1® reagent cartridges contain ethyl alcohol and guanidine salts which are hazardous materials. Solutions containing guanidine salts will generate toxic fumes when combined with bleach. Use appropriate care and wear appropriate protective clothing and eyewear when handling. Be careful not to expose face or hands to splashes. Dispose of EZ1® consumables in appropriate waste containers.

10 References

FBI Laboratory Safety Manual

DNA Procedures Manual

Qiagen[®]. *QIAcube*[®] *User Manual*, June 2008.

Qiagen[®]. EZ1[®] Advanced XL User Manual. May 2009.

Qiagen[®]. EZ1[®] DNA Investigator Handbook. April 2009.

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Date: 06/01/2021

Rev. #	Issue Date	History
3	02/15/19	2 & 4.5: Added Vacufuge contentrator in addition to Speed vac and
		default settings.
		4.1.4 & 4.3.5: Added allowance for additional spins and instruction
		to note any volume discrepancies after spins or elution.
		Appendix C: Changed QC samples to swabs.
4	06/01/21	Removed QIAsymphony throughout. Editorial changes throughout.
		1: Updated to SBAU.
		2: Added incubator/thermomisers. Added use info to Buffer ATL.
		3: Added instruction for RB to be the last sample of a batch. Added
		interpretation SOP reference numbers.
		4: Removed detail from any network computer.
		4.1 & 4.2: Relocated and added to L&S basket guidance.
		4.2.3: Relocated thermomixer note.
		4.2.5: A shaker rack plug must be in place next to each F fraction
		collection tube.
		4.2.9: Added guidance is excess supernatant needs to be removed.
		4.3.6: Added Vacufuge.
		4.4: Removed Speed-vac/Vacufuge procedures duplicated in quant
		SOP and added reference to DNA 226.
		App B, 3: Edited how minimum value established.

DNA Technical Leader Date: 06/01/2021 DCU Chief Date: 06/01/2021

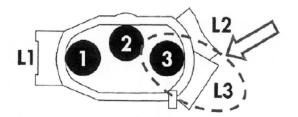
Redact - Signatures on File

Approval

SBAU Chief

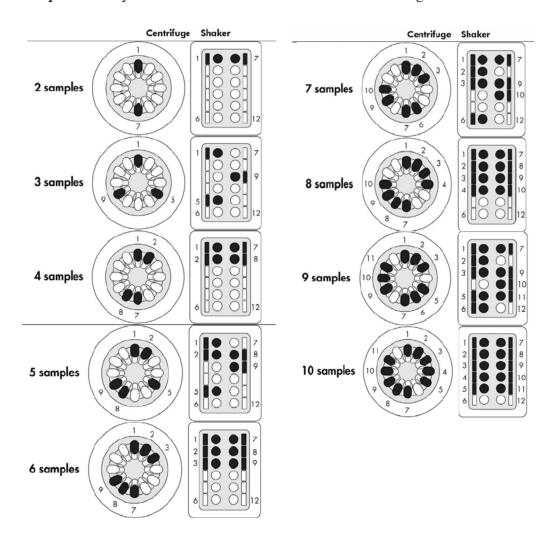
Appendix A: Loading the QIAcube®

The lysate tubes fit into position 3 of the rotor adaptors, with their caps folded back and inserted into position L3.





Rotor adaptors with lysate tubes must be distributed on the centrifuge for balance.



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Appendix A: Loading the QIAcube® (continued)

An overview of the loaded QIAcube®. (All caps must be removed prior to starting a run)



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Appendix B: Instrument General Maintenance and Performance Verification Procedures

Refer to the DNA procedure for instrument calibration and maintenance (i.e., DNAQA 608) for minimum frequency of performance verifications and additional requirements.

1. QIAcube® Robotic System (Extraction)

A. General Maintenance

There is no general maintenance required for the QIAcube[®].

B. Performance Verification

A portion of a swab containing 5 μ L female blood and 1 μ L of a 10x dilution of semen (semen diluted with reagent grade water) with known typing results will be processed using the QIAcube[®] differential lysis procedure followed by extraction using the appropriate DNA standard operating procedure (SOP).

The mixture must yield a distinguishable M fraction containing predominantly male DNA with minimal carryover from the F fraction as determined by quantification and amplification. The mixture is expected to yield a distinguishable F fraction containing predominantly female DNA but may contain a mixture of male and female DNA depending on the sample.

If the sample does not produce a distinguishable M fraction the process should be repeated with a new sample. If a predominately male DNA containing M fraction is not accomplished after 2 attempts the Technical Leader (TL) will be consulted.

2. EZ1® Advanced XL Robotic System (Extraction)

A. General Maintenance

Quarterly: Inspect the O-rings and grease as needed. Test Heating Block at 70°C.

B. Performance Verification

A swab containing 5 μ L blood from a known donor will be processed using the Normal lysis procedure followed by the EZ1® extraction procedure found in the appropriate DNA SOP for each of the 14 channels of the EZ1 robot. Alternately, individual channels may be tested to verify performance as needed.

A sample from each channel must yield a concentration, determined by quantification, greater than the minimum value established for the specific donor batch on the EZ1 $^{\text{@}}$ using a 50 μ L elution volume. A channel, or channels, may be repeated if necessary.

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If a channel, or channels, does not yield the appropriate concentration after 2 attempts the TL will be consulted.

3. Establishing the minimum value for Extraction Robot PV

Use an in service instrument and the appropriate extraction procedure to extract several samples from a blood card (i.e., FTA card) or swabs spotted with donor blood. The minimum quantitation value will be established by calculating 50% of the average of the quantitation values of the extracted samples.